

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/585,466
Source: 1 FWP
Date Processed by STIC: 7/21/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 07/21/2006

PATENT APPLICATION: US/10/585,466

TIME: 18:33:54

Input Set : N:\Crf4\07192006\J585466.raw

Output Set: N:\CRF4\07212006\J585466.raw

1 <110> APPLICANT: Avalon Pharmaceuticals

2 <120> TITLE OF INVENTION: Cancer-Linked Genes as Targets for Chemotherapy

3 <130> FILE REFERENCE: 689290-272

4 <140> CURRENT APPLICATION NUMBER: US/10/585,466

5 <141> CURRENT FILING DATE: 2006-07-05

6 <150> PRIOR APPLICATION NUMBER: 60/534,419

7 <151> PRIOR FILING DATE: 2004-01-06

8 <160> NUMBER OF SEQ ID NOS: 69

9 <170> SOFTWARE: PatentIn version 3.3

11 <210> SEQ ID NO: 1

12 <211> LENGTH: 1181

13 <212> TYPE: DNA

14 <213> ORGANISM: Artificial

15 <220> FEATURE:

16 <223> OTHER INFORMATION: cDNA or putative protein derived from a cDNA.

17 <400> SEQUENCE: 1

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21	gcggccgctc cctcgtctcg tccccagact catggggcag caccccagcc gacagccccg	240
22	tggcgagccc cgcgcggcca ggcaccctcc gggacccccg cgccccctcc gtaggtaggc	300
23	gcggcgcgcg cagcagccgc ctgggcagcg ggcagaggca gagcgccagt gagcgggaga	360
24	aactgcgcat gcgcacgctg gcccgcgccc tgcacgagct gcgccgcttt ctaccgccgt	420
25	ccgtggcgcc cgcggggccag agcctgacca agatcgagac gctgcgcctg gctatccgct	480
26	atatcggcca cctgtcggcc gtgctaggcc tcagcgagga gagtctccag cgccggtgcc	540
27	ggcagcgcgg tgacgcgggg tcccctcggg gctgcccgct gtgccccgac gactgccccg	600
28	cgcagatgca gacacggacg caggctgagg ggcaggggca ggggcgcggg ctgggcctgg	660
29	tatccgccgt ccgcgcgggg gcgtcctggg gatccccgcc tgctgcccc ggagcccag	720
30	ctgcacccga gccgcgcgac ccgcctgcgc tgttcgccga ggcggcgtgc ccggaagggc	780
31	aggcgatgga gccaaagcca ccgtccccgc tccttcggg cgacgtgctg gctctgttgg	840
32	agacctggat gccctctcgc cctctggagt ggctgcctga ggagcccaag tgacaaggga	900
33	caactgacgc cgtctctgtg agcaccgagg ctttttggcc tcagcacctt cgaagtgggt	960
34	ccttggcaga ctgcctttcc tggaaagagg caccggcgat cccgacgggg gcattcctgc	1020
35	gggtgagagc cgtccccacc gcggcgggccc ttctcagccc ctccctccat ggagggaccc	1080
36	atagggctag acactttgag gcaagcagga ggctctgcct aatgtgaatt tatttatttg	1140
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40 <211> LENGTH: 268

41 <212> TYPE: PRT

42 <213> ORGANISM: Artificial

43 <220> FEATURE:

44 <223> OTHER INFORMATION: cDNA or putative protein derived from a cDNA.

45 <400> SEQUENCE: 2

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47 1 5 10 15
48 Ala Ala Trp Gly Pro Thr Arg Arg Pro Pro Ser Asp Lys Asp Cys
49 20 25 30
50 Gly Arg Ser Leu Val Ser Ser Pro Asp Ser Trp Gly Ser Thr Pro Ala
51 35 40 45
52 Asp Ser Pro Val Ala Ser Pro Ala Arg Pro Gly Thr Leu Arg Asp Pro
53 50 55 60
54 Arg Ala Pro Ser Val Gly Arg Arg Gly Ala Arg Ser Ser Arg Leu Gly
55 65 70 75 80
56 Ser Gly Gln Arg Gln Ser Ala Ser Glu Arg Glu Lys Leu Arg Met Arg
57 85 90 95
58 Thr Leu Ala Arg Ala Leu His Glu Leu Arg Arg Phe Leu Pro Pro Ser
59 100 105 110
60 Val Ala Pro Ala Gly Gln Ser Leu Thr Lys Ile Glu Thr Leu Arg Leu
61 115 120 125
62 Ala Ile Arg Tyr Ile Gly His Leu Ser Ala Val Leu Gly Leu Ser Glu
63 130 135 140
64 Glu Ser Leu Gln Arg Arg Cys Arg Gln Arg Gly Asp Ala Gly Ser Pro
65 145 150 155 160
66 Arg Gly Cys Pro Leu Cys Pro Asp Asp Cys Pro Ala Gln Met Gln Thr
67 165 170 175
68 Arg Thr Gln Ala Glu Gly Gln Gly Gln Gly Arg Gly Leu Gly Leu Val
69 180 185 190
70 Ser Ala Val Arg Ala Gly Ala Ser Trp Gly Ser Pro Pro Ala Cys Pro
71 195 200 205
72 Gly Ala Arg Ala Ala Pro Glu Pro Arg Asp Pro Pro Ala Leu Phe Ala
73 210 215 220
74 Glu Ala Ala Cys Pro Glu Gly Gln Ala Met Glu Pro Ser Pro Pro Ser
75 225 230 235 240
76 Pro Leu Leu Pro Gly Asp Val Leu Ala Leu Leu Glu Thr Trp Met Pro
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90 tctgcctggc tgctctgtgc tctgtgtcct ctcctttctt tcgcttctc caaacattgc 180
91 tccttcaatc ccaggagaag tctcctcgga tgtcagcgcc tctaaagcag cccaaggctt 240
92 gcctcaattg catggtttcc cgagtcctca gctccagaag accaggcaga tgggtggacc 300
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94 ggcagcgagg aggaggaggt ttctccaagg acagaag 397
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105     ccctcccggc ttcttttctt tgtagccacc tcagggggaag caacagatcg tcaactcggtg    180
106     ttctcaccga aagcacgtaa tcgcccgtgt aactcatggt ggctgggggg cctcccgggc    240
107     cgcgcggaga ggctgggggt cgcctccatg cagcatgctt gtgctcaatt gcagggtcct    300
108     cggttctcag tgtgcagagg gcggtgagag ctcaactctc gtcccccact cccaccgcga    360
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110     gtgtggcctg tgggtctcga gtctgttctc ctggagtctc gcatttgcac ccccttcttc    480
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113     cacggtgact tgattacact ctctcattca tggtcacttc cgaagcgctt tagtgcttct    660
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115     ggcccggaag ggccttcgtc ttaccgggga tccacctctc cctcatctt ccctgcctac    780
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144     Cys Phe Ser Lys Leu Gln Gly Pro Gly Ser Pro Leu Gln Gly Arg Ala
145           35           40           45
146     Pro Leu Leu Pro Ala Arg His Trp Arg Trp Pro Pro Glu Arg Arg Arg

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154 <213> ORGANISM: Artificial
155 <220> FEATURE:
156 <223> OTHER INFORMATION: cDNA or putative protein derived from a cDNA.
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160      ccctcccggc ttcttttctt tgtagccacc tcaggggaag caacagatcg tcaactcggtg      180
161      ttctcaccga aagcacgtaa tcgcccgtgt aactcatggt ggctgggggg cctcccggcg      240
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163      cgttctcgag tgtgcagagg gcggtgagag ctcaactctc gtccccacct cccaccgcga      360
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191      cctaccttcg aacaaaagga atgcatgaag ggtttcagtg actttgccat aacaaaggcg      2040
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193      tagggctttt aagaccataa ttgaaccggg gggcgtctag gaaaaccgaa aacagttcta      2160
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231      Pro Gln Glu Thr Lys Ala Asn Ser Gly Glu Lys Ala Gly Ala Glu Lys
232      35          40          45
233      Gly Ala Pro Gly Ala Ala Gly Arg Ala Ile Arg Gln Ala Ala Pro Lys
234      50          55          60
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236      65          70          75          80
237      Leu Pro
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:20; N Pos. 2,6,14,725,782,828

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

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Seq#:52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69

VERIFICATION SUMMARY

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L:773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0

M:341 Repeated in SeqNo=20